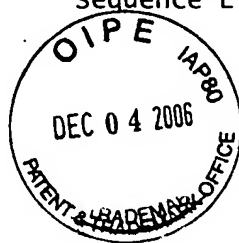


<110> Genentech Inc.
Ashkenazi, Avi
Fong, Sherman
Goddard, Audrey
Gurney, Austin L.
Napier, Mary A.
Tumas, Daniel
Wood, William I.



<120> COMPOUNDS, COMPOSITIONS AND METHODS FOR THE TREATMENT
OF DISEASES CHARACTERIZED BY A-33 RELATED ANTIGENS

<130> P1216R1

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<141> 1998-11-20

<150> US 60/066,364

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Glu Trp Lys Phe Asp Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr
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Ile Ser Val Pro Ser Ser Val Thr Ile Gly Asn Arg Ala Val Leu
Page 10

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 35 40 45
 Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr Ala Ser Tyr
 50 55 60
 Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr Phe Lys Ser
 65 70 75
 Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met Val Ser Glu
 80 85 90
 Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys Leu Ile Val
 95 100 105
 Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro Ser Ser Ala
 110 115 120
 Thr Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu Gln Asp Gly
 125 130 135
 Ser Pro Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly Ile Val Met
 140 145 150
 Pro Thr Asn Pro Lys Ser Thr Arg Ala Phe Ser Asn Ser Ser Tyr
 155 160 165
 Val Leu Asn Pro Thr Thr Gly Glu Leu Val Phe Asp Pro Leu Ser
 170 175 180
 Ala Ser Asp Thr Gly Glu Tyr Ser Cys Glu Ala Arg Asn Gly Tyr
 185 190 195
 Gly Thr Pro Met Thr Ser Asn Ala Val Arg Met Glu Ala Val Glu
 200 205 210
 Arg Asn Val Gly Val Ile Val Ala Ala Val Leu Val Thr Leu Ile
 215 220 225
 Leu Leu Gly Ile Leu Val Phe Gly Ile Trp Phe Ala Tyr Ser Arg
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 245 250 255
 Ile Tyr Ser Gln Pro
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          35          40          45
Lys Leu Leu Leu Thr His Thr Glu Arg Val Val Ile Trp Pro Phe
          50          55          60
Ser Asn Lys Asn Tyr Ile His Gly Glu Leu Tyr Lys Asn Arg Val
          65          70          75
Ser Ile Ser Asn Asn Ala Glu Gln Ser Asp Ala Ser Ile Thr Ile
          80          85          90
Asp Gln Leu Thr Met Ala Asp Asn Gly Thr Tyr Glu Cys Ser Val
          95          100          105
Ser Leu Met Ser Asp Leu Glu Gly Asn Thr Lys Ser Arg Val Arg
          110          115          120
Leu Leu Val Leu Val Pro Pro Ser Lys Pro Glu Cys Gly Ile Glu
          125          130          135
Gly Glu Thr Ile Ile Gly Asn Asn Ile Gln Leu Thr Cys Gln Ser
          140          145          150
Lys Glu Gly Ser Pro Thr Pro Gln Tyr Ser Trp Lys Arg Tyr Asn
          155          160          165
Ile Leu Asn Gln Glu Gln Pro Leu Ala Gln Pro Ala Ser Gly Gln
          170          175          180
Pro Val Ser Leu Lys Asn Ile Ser Thr Asp Thr Ser Gly Tyr Tyr
          185          190          195
Ile Cys Thr Ser Ser Asn Glu Glu Gly Thr Gln Phe Cys Asn Ile
          200          205          210
Thr Val Ala Val Arg Ser Pro Ser Met Asn Val Ala Leu Tyr Val
          215          220          225
Gly Ile Ala Val Gly Val Val Ala Ala Leu Ile Ile Ile Gly Ile
          230          235          240
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 35 40 45

Gln Gly Asp Thr Thr Arg Leu Val Cys Tyr Asn Asn Lys Ile Thr
 50 55 60

Ala Ser Tyr Glu Asp Arg Val Thr Phe Leu Pro Thr Gly Ile Thr
 65 70 75

Phe Lys Ser Val Thr Arg Glu Asp Thr Gly Thr Tyr Thr Cys Met
 80 85 90

Val Ser Glu Glu Gly Gly Asn Ser Tyr Gly Glu Val Lys Val Lys
 95 100 105

Leu Ile Val Leu Val Pro Pro Ser Lys Pro Thr Val Asn Ile Pro
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Ser Ser Ala Thr Ile Gly Asn Arg Ala Val Leu Thr Cys Ser Glu
 125 130 135

Gln Asp Gly Ser Pro Pro Ser Glu Tyr Thr Trp Phe Lys Asp Gly
 140 145 150

Ile Val Met Pro Thr Asn Pro Lys Ser Thr Arg Ala Phe Ser Asn
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Ser Ser Tyr Val Leu Asn Pro Thr Thr Gly Glu Leu Val Phe Asp
 170 175 180

Pro Leu Ser Ala Ser Asp Thr Gly Glu Tyr Ser Cys Glu Ala Arg
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Asn Gly Tyr Gly Thr Pro Met Thr Ser Asn Ala Val Arg Met Glu
 200 205 210

Ala Val Glu Arg Asn Val Gly Val Ile Val Ala Ala Val Leu Val
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Thr Leu Ile Leu Leu Gly Ile Leu Val Phe Gly Ile Trp Phe Ala
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Lys Lys Val Ile Tyr Ser Gln Pro
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<213> Homo sapiens

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Leu Cys Ala Val Arg Val Thr Val Asp Ala Ile Ser Val Glu Thr
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1

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10

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Pro Cys Thr Tyr His Thr Ser Thr Ser Ser Arg Glu Gly Leu Ile
35 40 45

Gln Trp Asp Lys Leu Leu Leu Thr His Thr Glu Arg Val Val Ile
50 55 60

Trp Pro Phe Ser Asn Lys Asn Tyr Ile His Gly Glu Leu Tyr Lys
65 70 75

Asn Arg Val Ser Ile Ser Asn Asn Ala Glu Gln Ser Asp Ala Ser
80 85 90

Ile Thr Ile Asp Gln Leu Thr Met Ala Asp Asn Gly Thr Tyr Glu
95 100 105

Cys Ser Val Ser Leu Met Ser Asp Leu Glu Gly Asn Thr Lys Ser
110 115 120

Arg Val Arg Leu Leu Val Leu Val Pro Pro Ser Lys Pro Glu Cys
125 130 135

Gly Ile Glu Gly Glu Thr Ile Ile Gly Asn Asn Ile Gln Leu Thr
140 145 150

Cys Gln Ser Lys Glu Gly Ser Pro Thr Pro Gln Tyr Ser Trp Lys
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Arg Tyr Asn Ile Leu Asn Gln Glu Gln Pro Leu Ala Gln Pro Ala
170 175 180

Ser Gly Gln Pro Val Ser Leu Lys Asn Ile Ser Thr Asp Thr Ser
185 190 195

Gly Tyr Tyr Ile Cys Thr Ser Ser Asn Glu Glu Gly Thr Gln Phe
200 205 210

Cys Asn Ile Thr Val Ala Val Arg Ser Pro Ser Met Asn Val Ala
215 220 225

Leu Tyr Val Gly Ile Ala Val Gly Val Val Ala Ala Leu Ile Ile
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cttggtccc aaagcacca cagctcatac acaatgaata caaaaactgg 300
aactctgcaa ttaatactg ttccaaact ggacactgga gaatattcct 350
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